

“Servicios ecosistémicos, ecología de las enfermedades y bienestar humano”

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Oikos, Capacitación para la sostenibilidad

**Lanzamiento del Fondo Internacional para Proyectos de
Investigación en Ecosistemas (ESPA)**

Miraflores, 26 de Enero del 2010



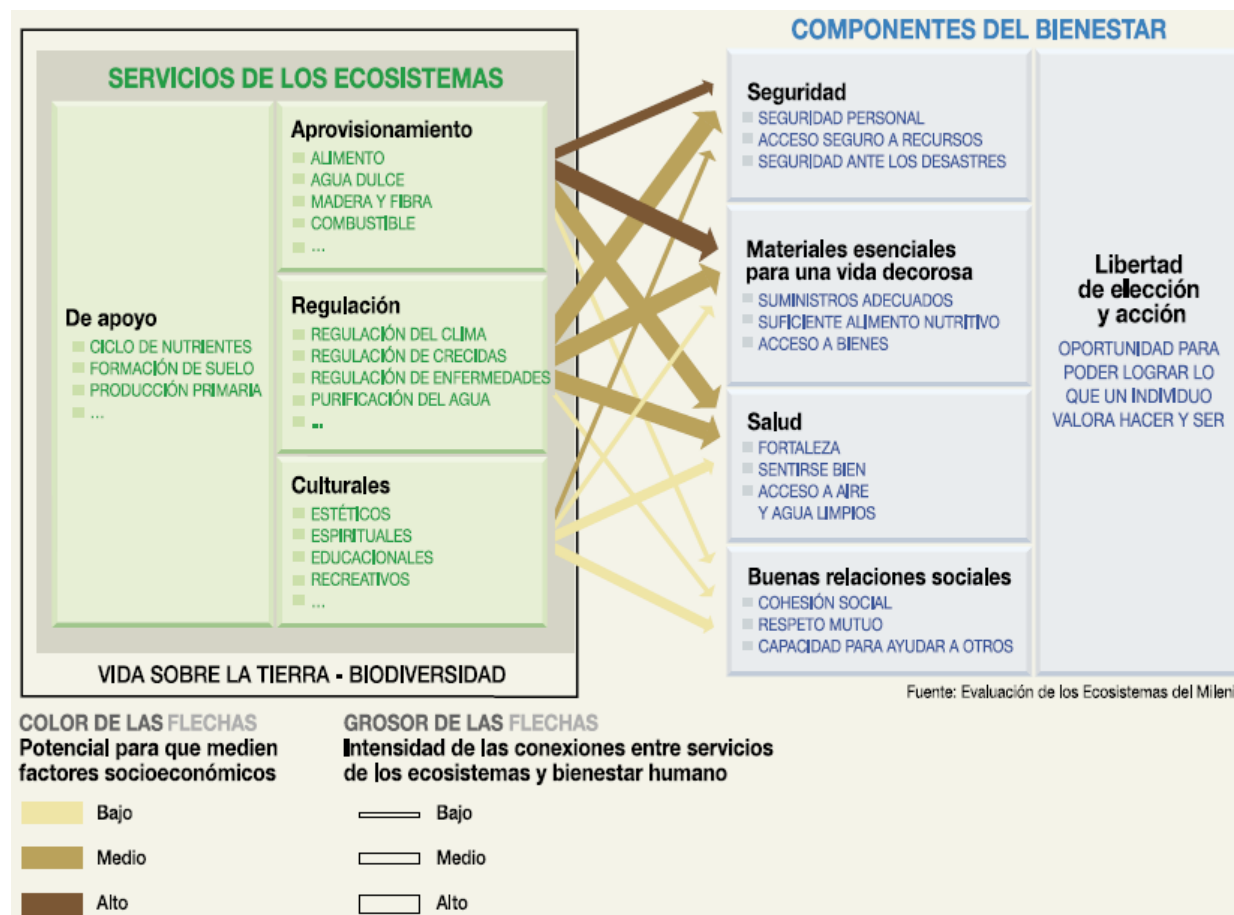
Servicios de los Ecosistemas



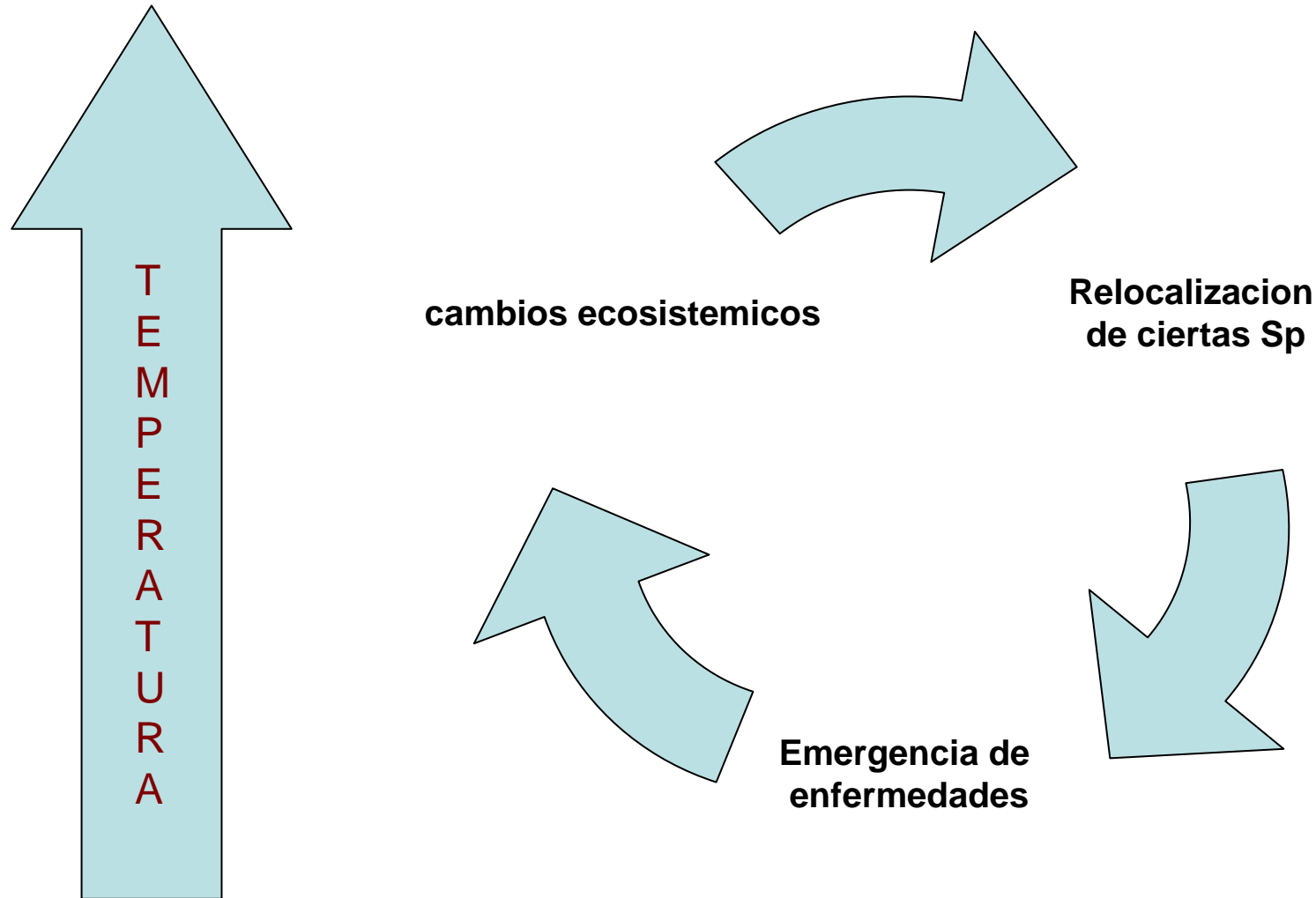
Los beneficios que la gente obtiene de los ecosistemas



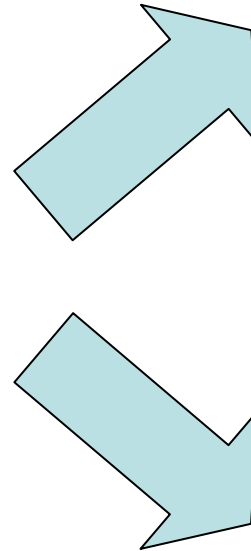
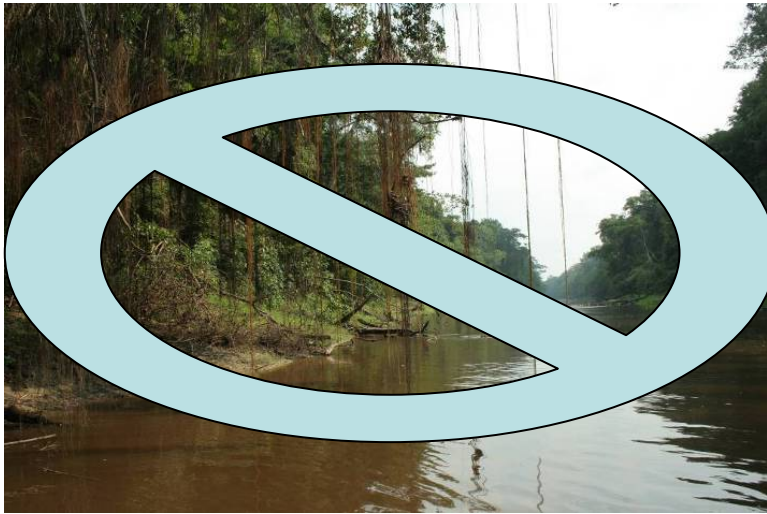
Consecuencias de los cambios en los ecosistemas para el bienestar humano

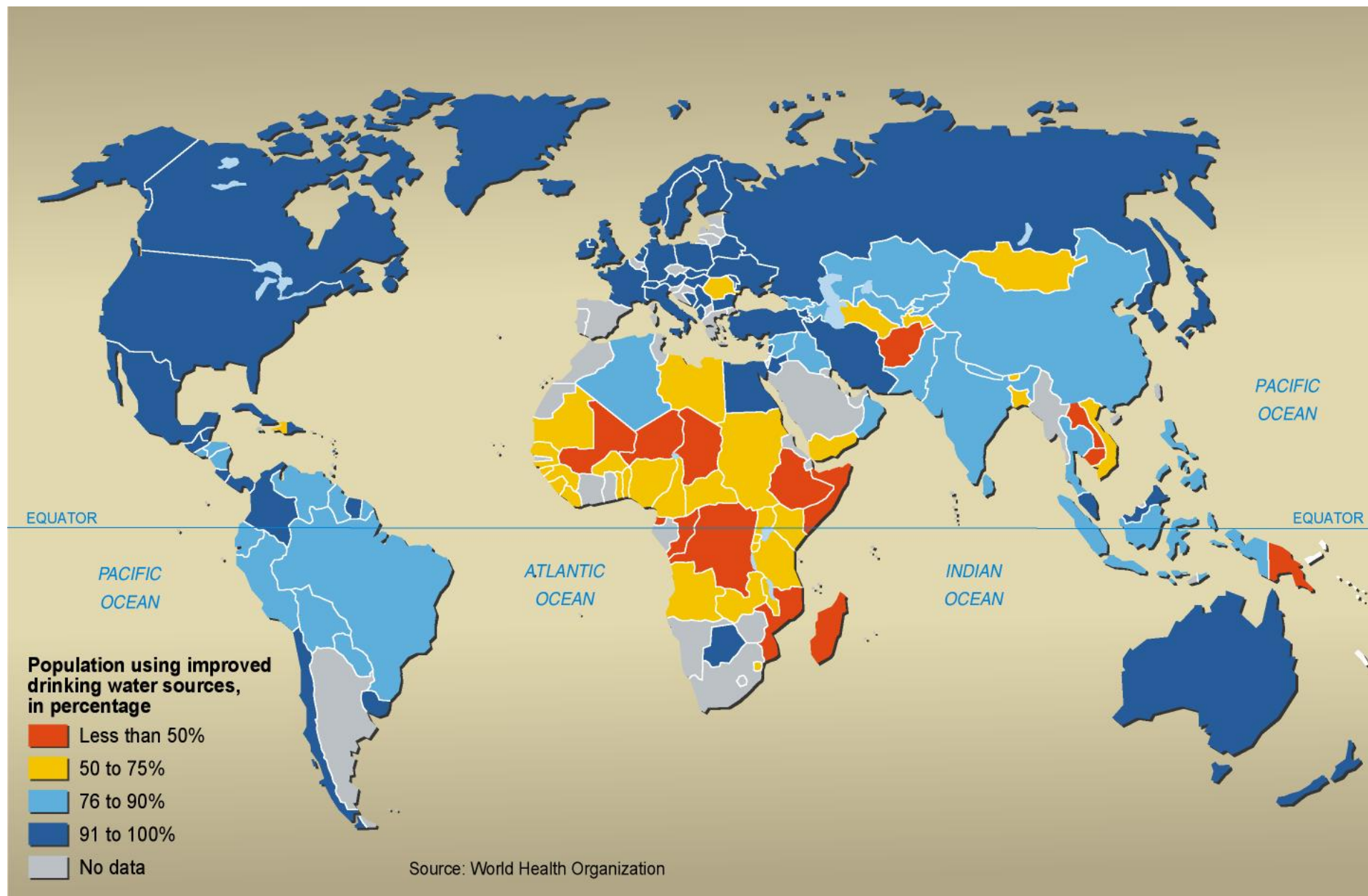


Regulación del clima

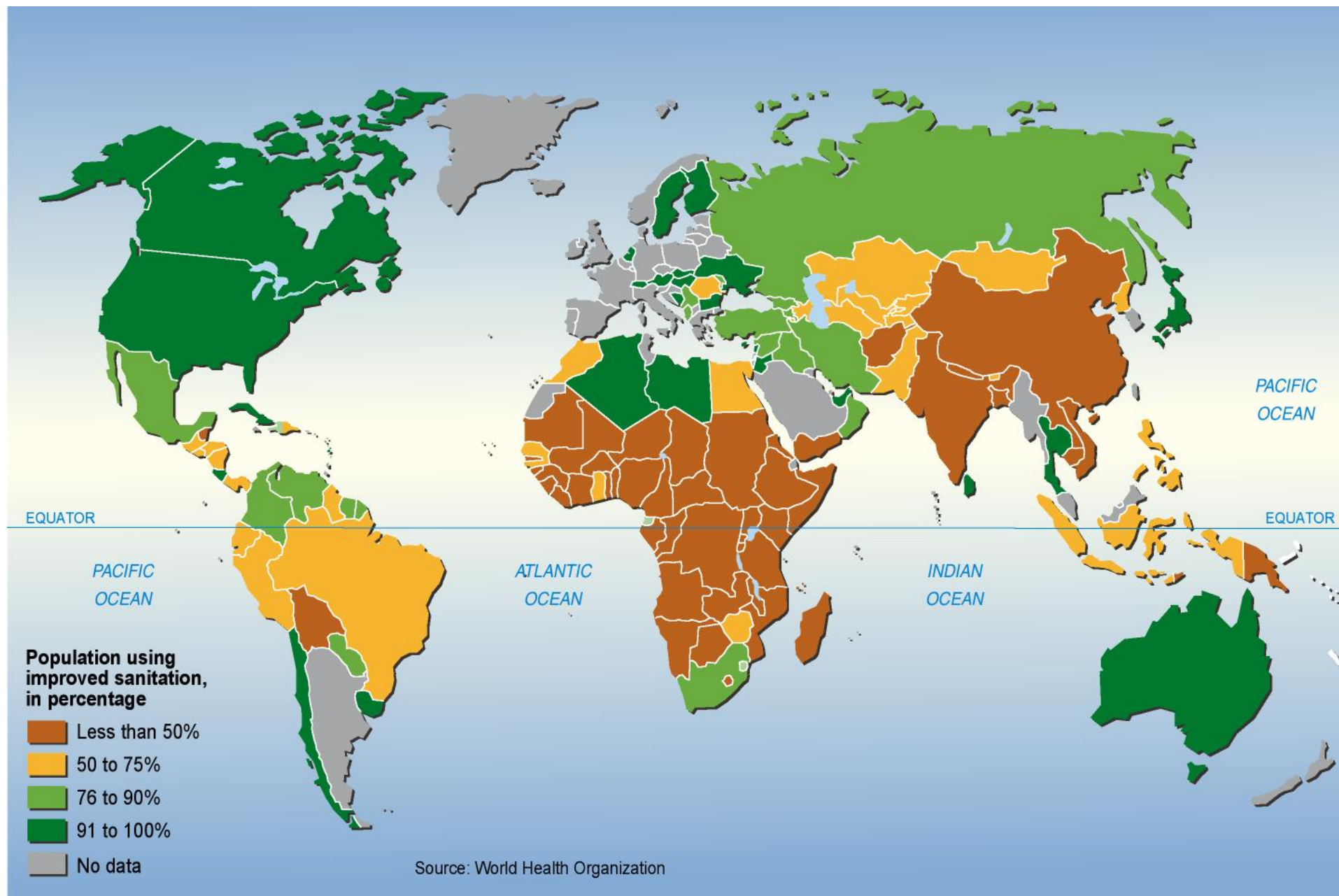


Purificación del agua





- Unos 1,100 millones de personas aún carecen de acceso a un suministro adecuado de agua
- La escasez de agua afecta entre 1,000 y 2,000 millones de personas en el mundo



- Más de 2,600 millones no tienen acceso a saneamiento adecuado

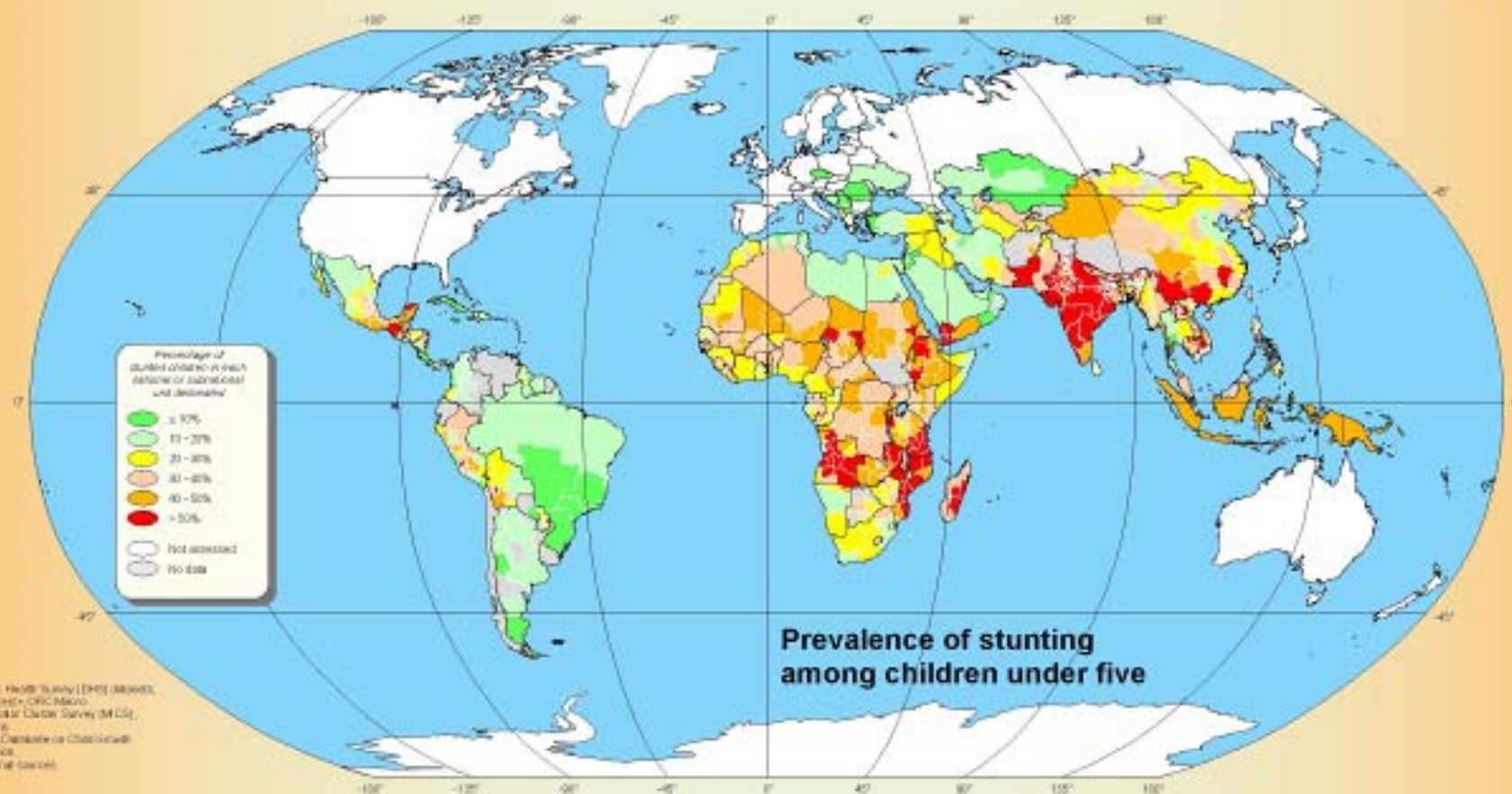


www.povertymap.net

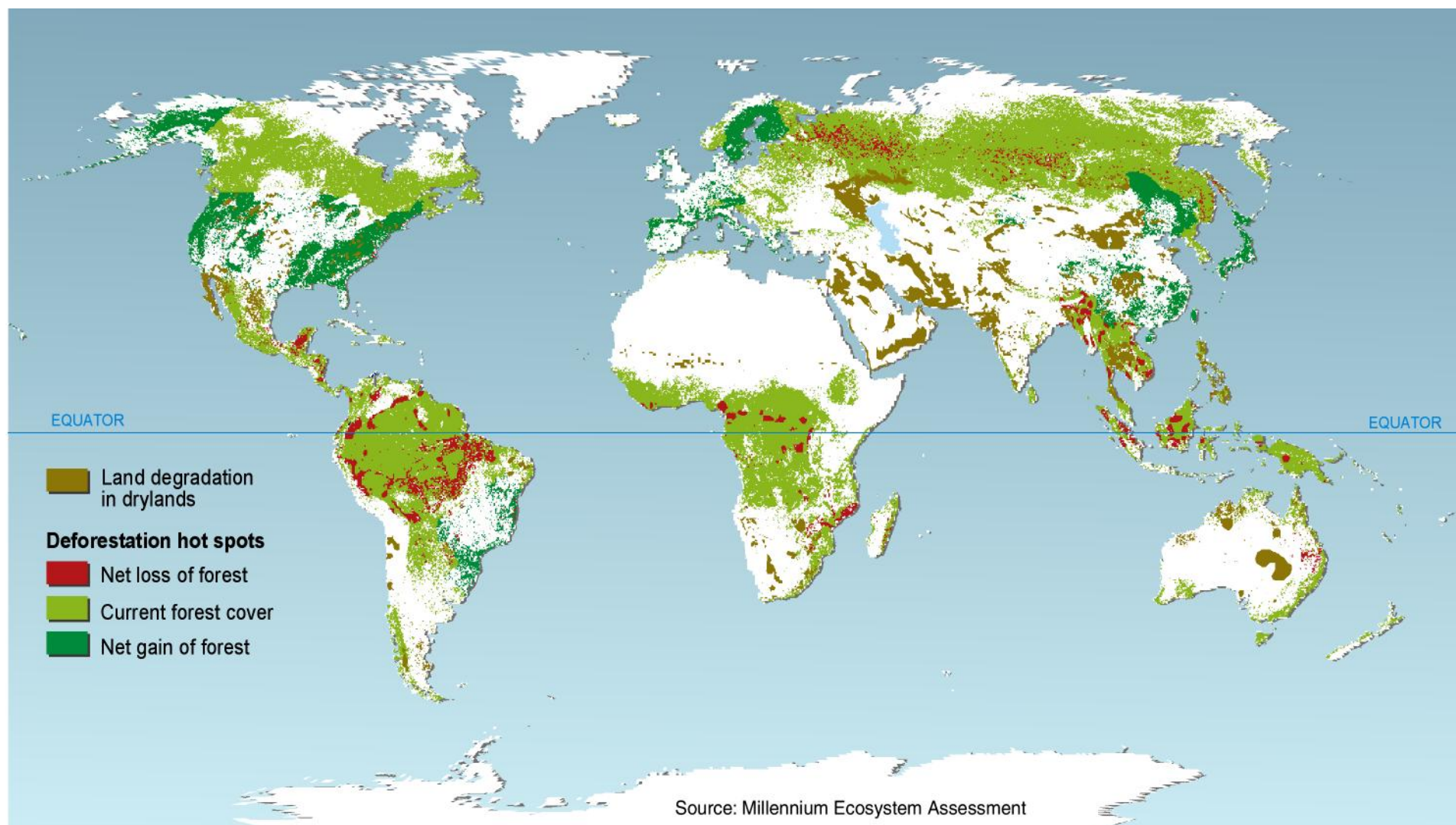
Chronic Undernutrition among Children An Indicator of Poverty



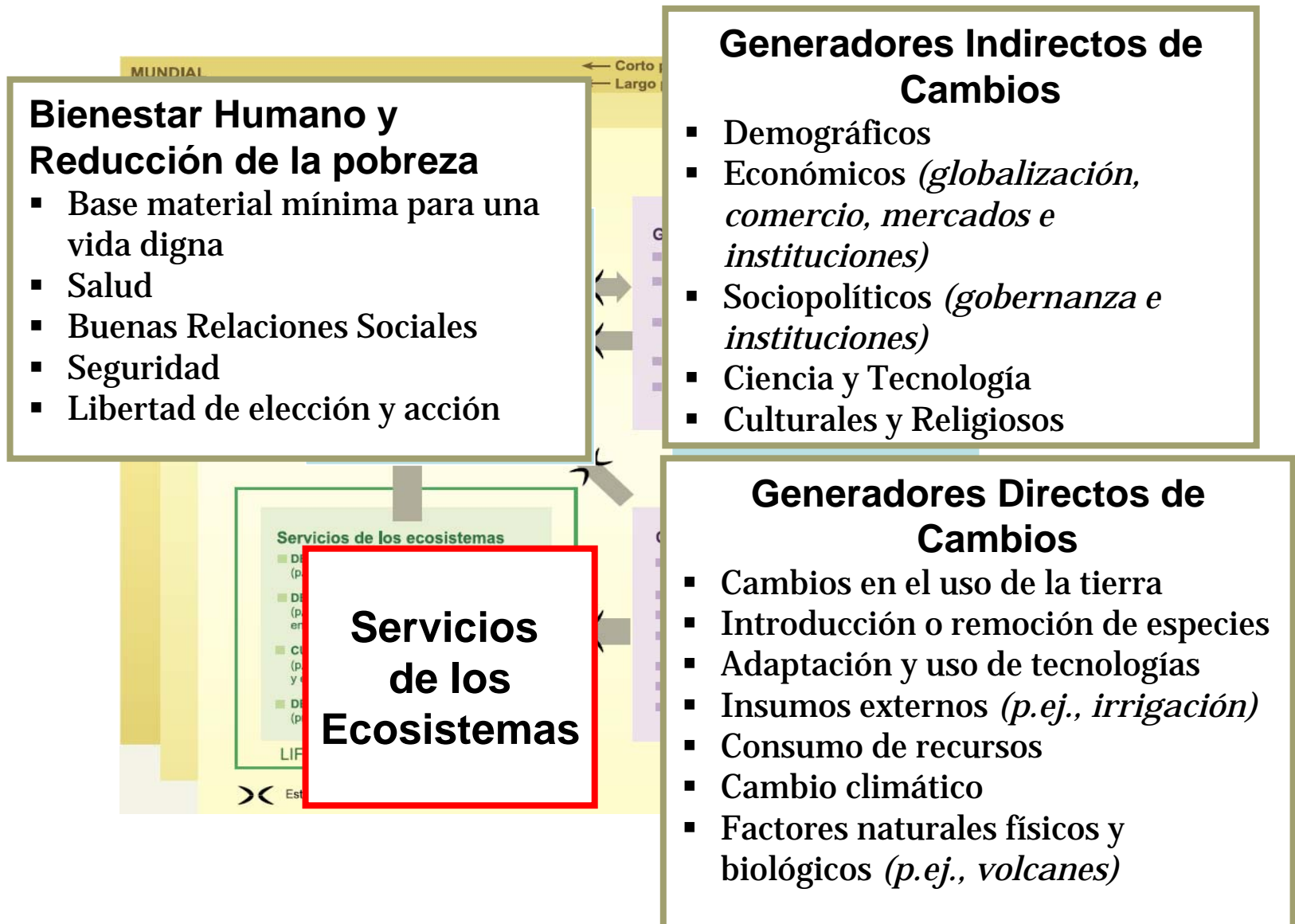
www.fao.org



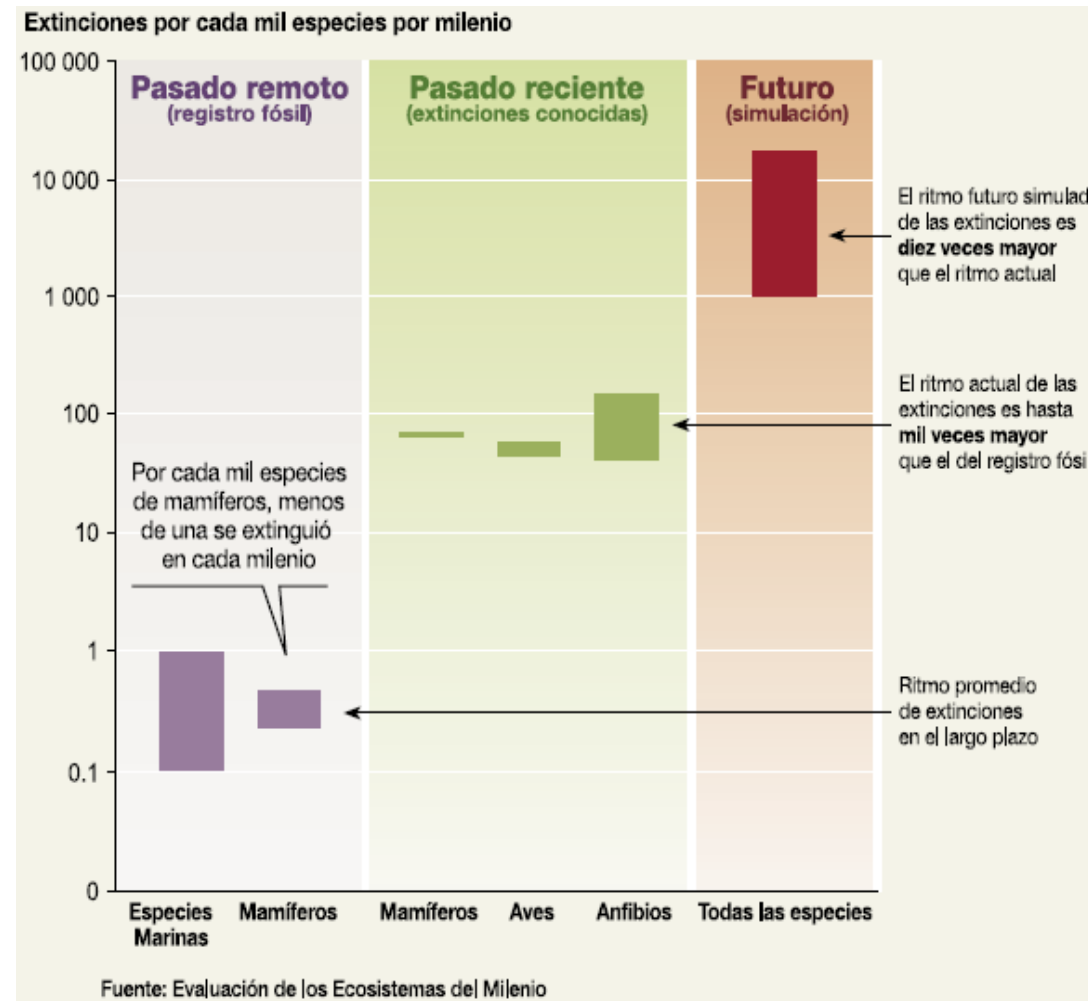
- Unas 852 millones de personas estaban desnutridas en el año 2000–02, 37 millones más que en el período 1997–99

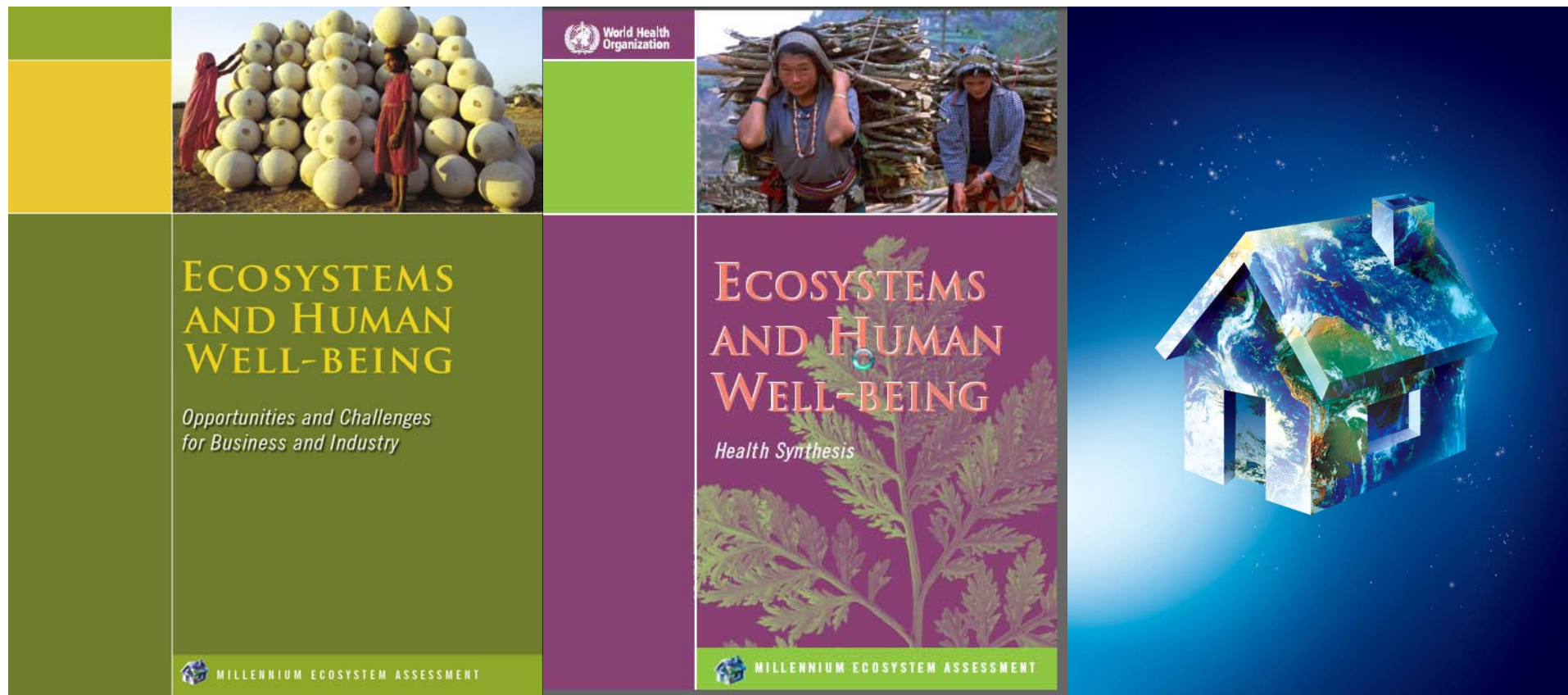


Marco conceptual de la EM



Cambios significativos y en gran medida irreversibles en la diversidad de las especies

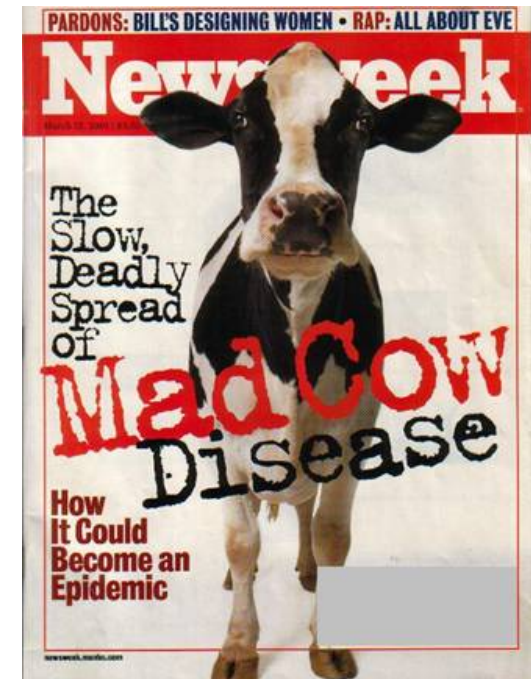




ENFERMEDADES EMERGENTES

Enfermedades que han aumentado su incidencia, han expandido su ámbito geográfico, han sido recientemente reconocidas o que presentan manifestaciones nuevas.

(Daszak et al. 2000a)



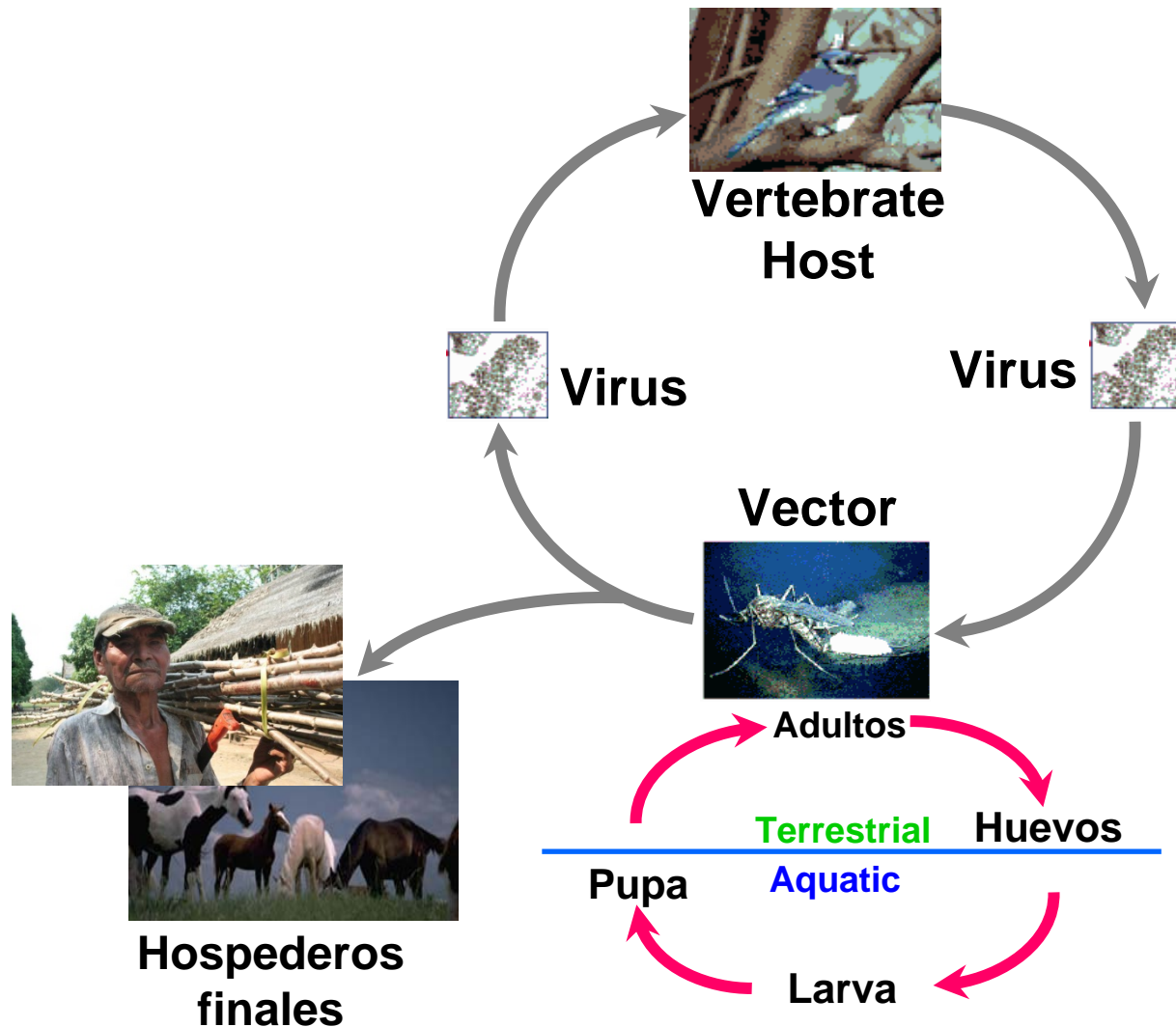
Ecología de enfermedades

- Nuevo paradigma:
 - Mejor entendimiento de la dinámica de las enfermedades teniendo en cuenta el componente ambiental.
 - Multicausalidad en el origen de las enfermedades genera la necesidad de un enfoque transdisciplinario para enfrentarlas.

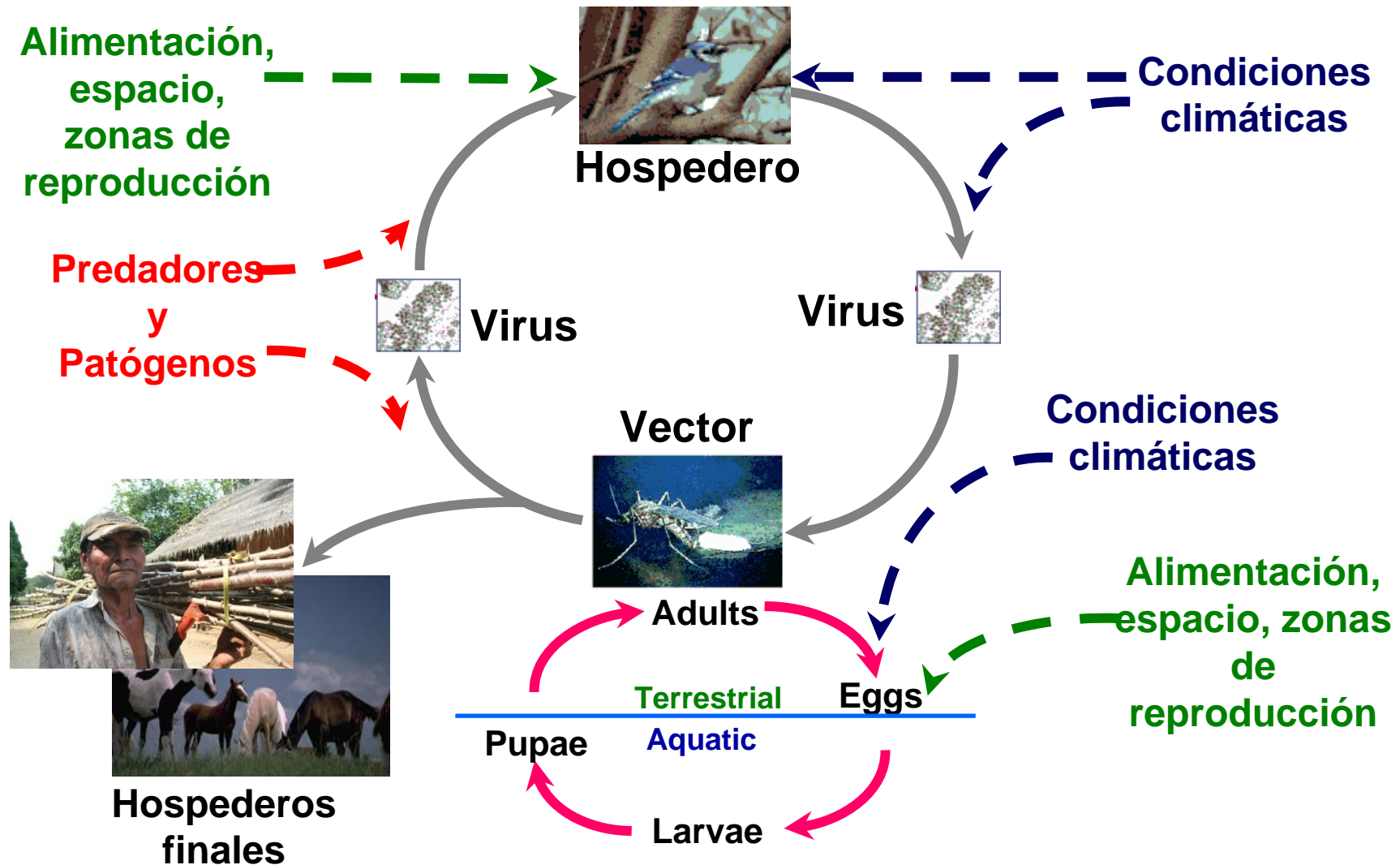


Arbovirus: Ciclo de Transmisión

(arthropod-borne viruses)



Arbovirus: Ciclo de Transmisión



LETTERS

Global trends in emerging infectious diseases

Kate E. Jones¹, Nikkita G. Patel², Marc A. Levy³, Adam Storeygard^{3†}, Deborah Balk^{3†}, John L. Gittleman⁴ & Peter Daszak²

- Las enfermedades emergentes son en su mayoría causadas por zoonosis (60.3%)
- Un 70.3% de las enfermedades emergentes causadas por zoonosis tienen origen en la fauna silvestre y tienen un incremento significativo a través del tiempo.



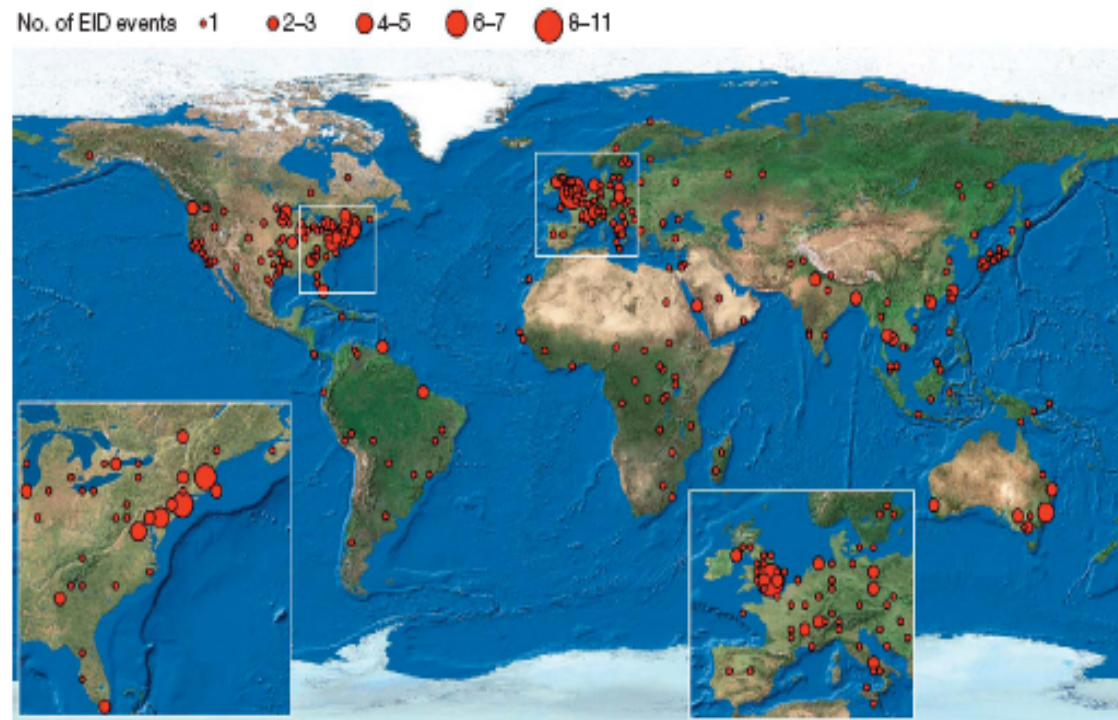
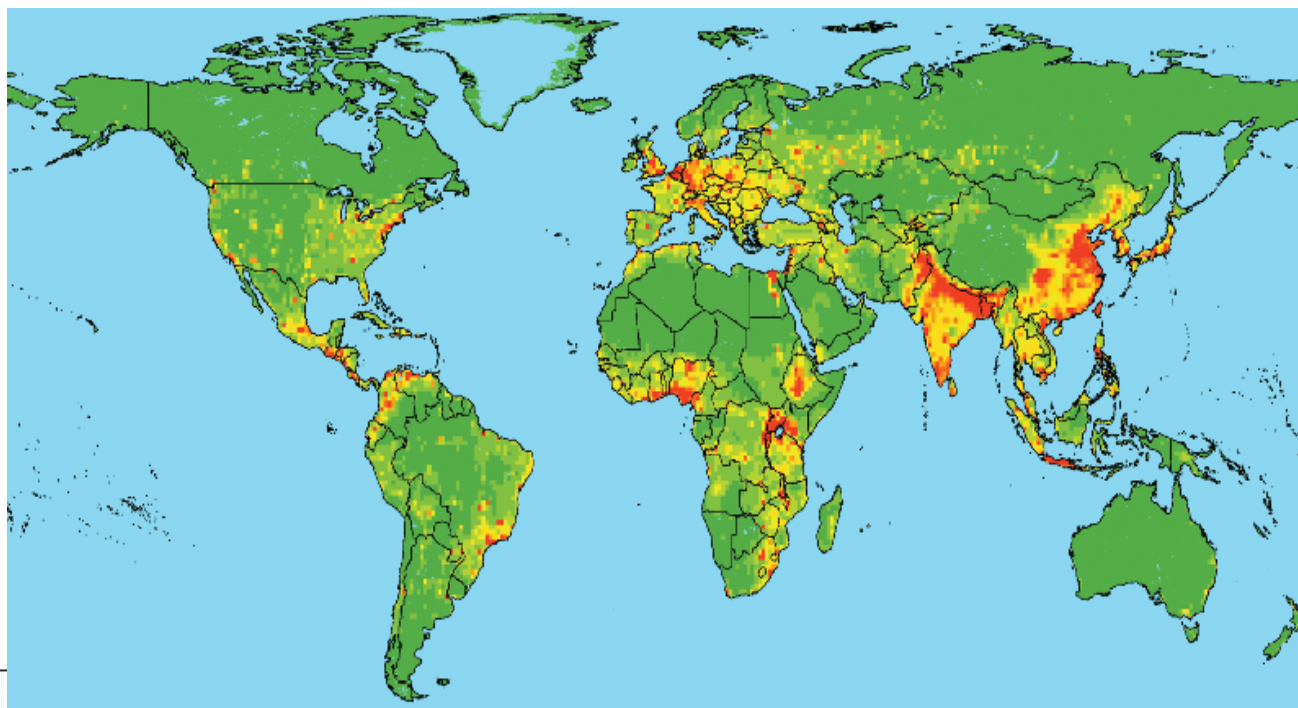
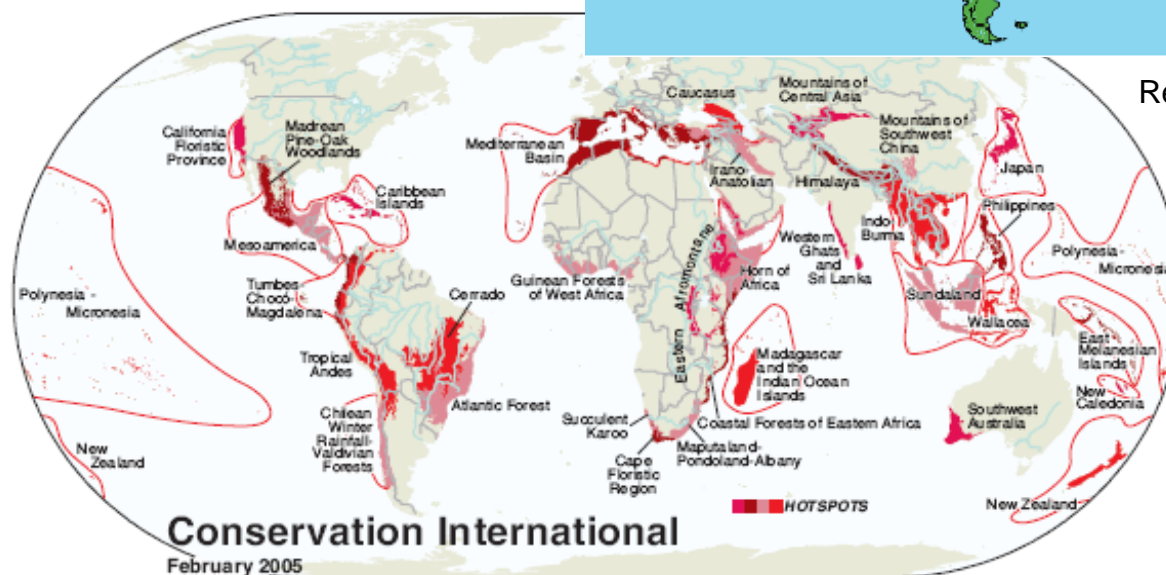


Figure 2 | Global richness map of the geographic origins of EID events from 1940 to 2004. The map is derived for EID events caused by all pathogen types. Circles represent one degree grid cells, and the area of the circle is proportional to the number of events in the cell.



BIODIVERSITY HOTSPOTS



Relative geographic risk of emerging or re-emerging infectious diseases that originate with zoonotic pathogens from wildlife



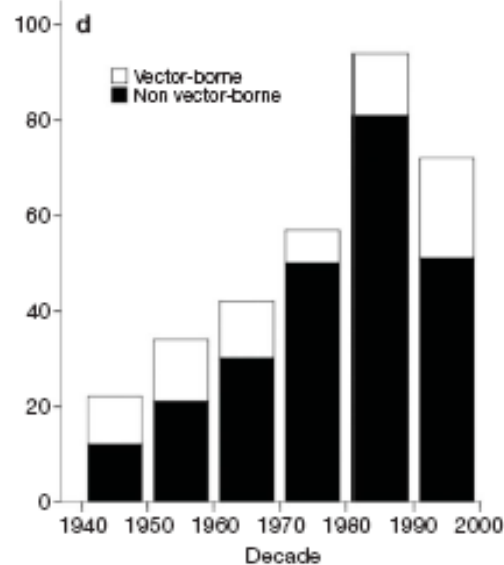
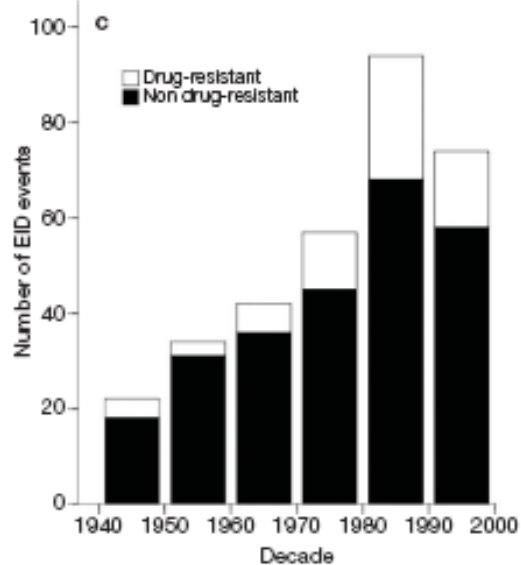
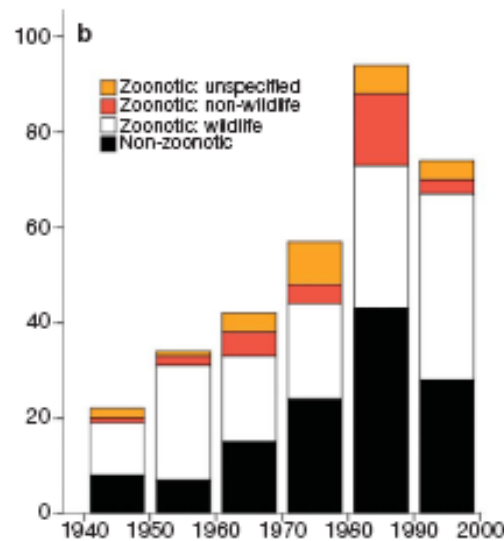
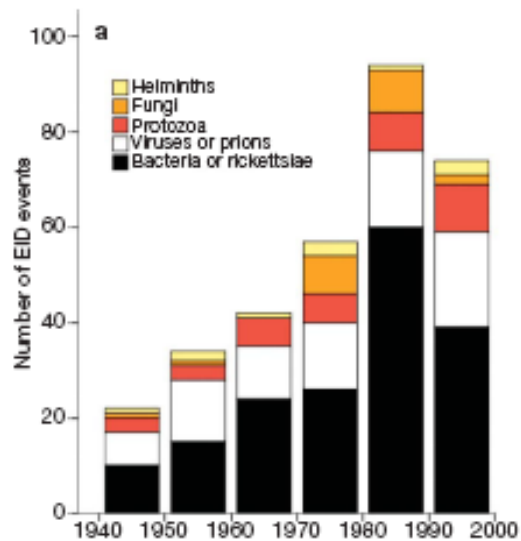
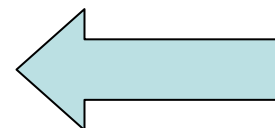
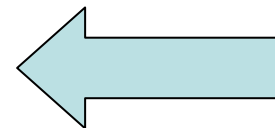
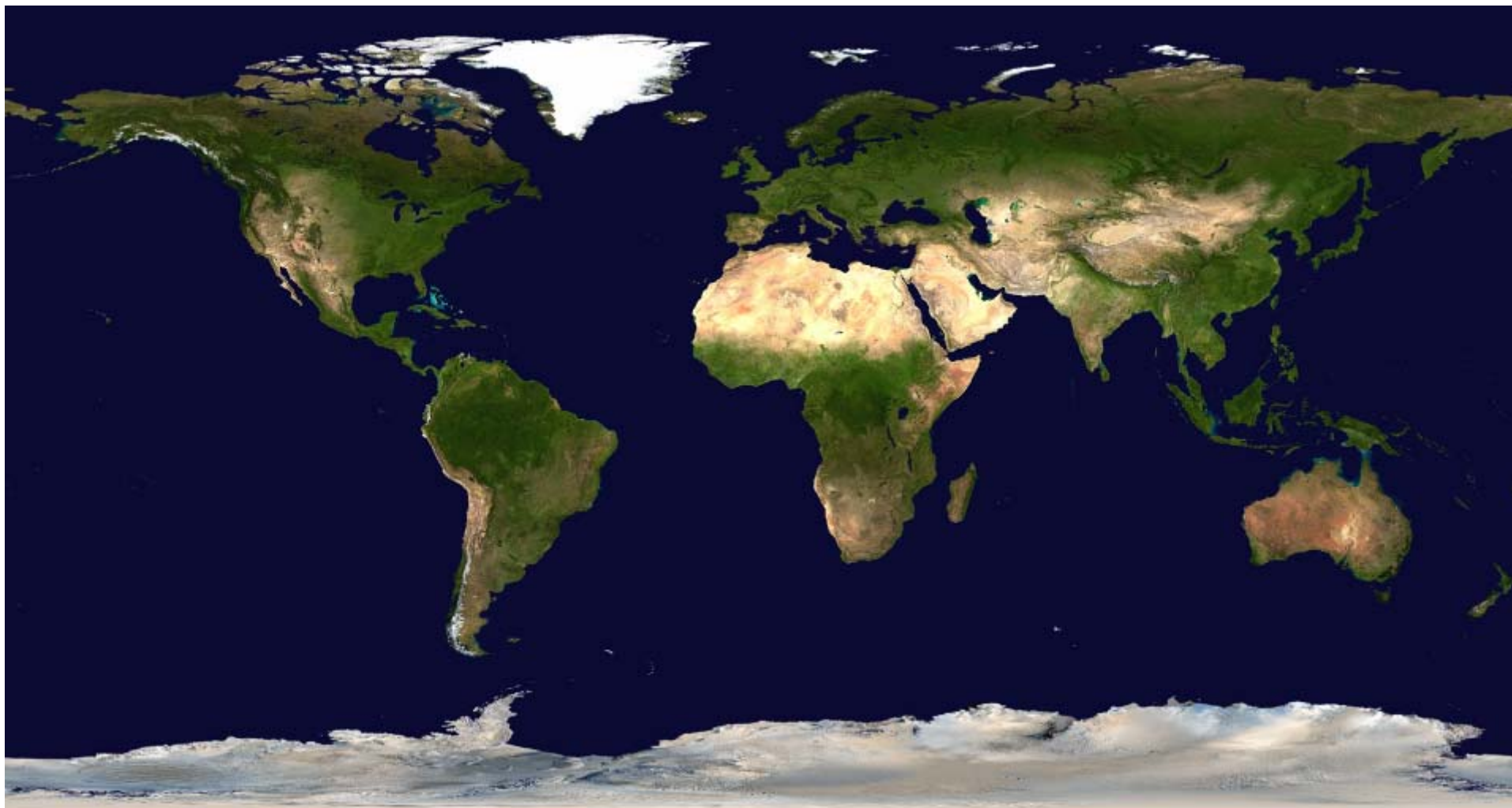


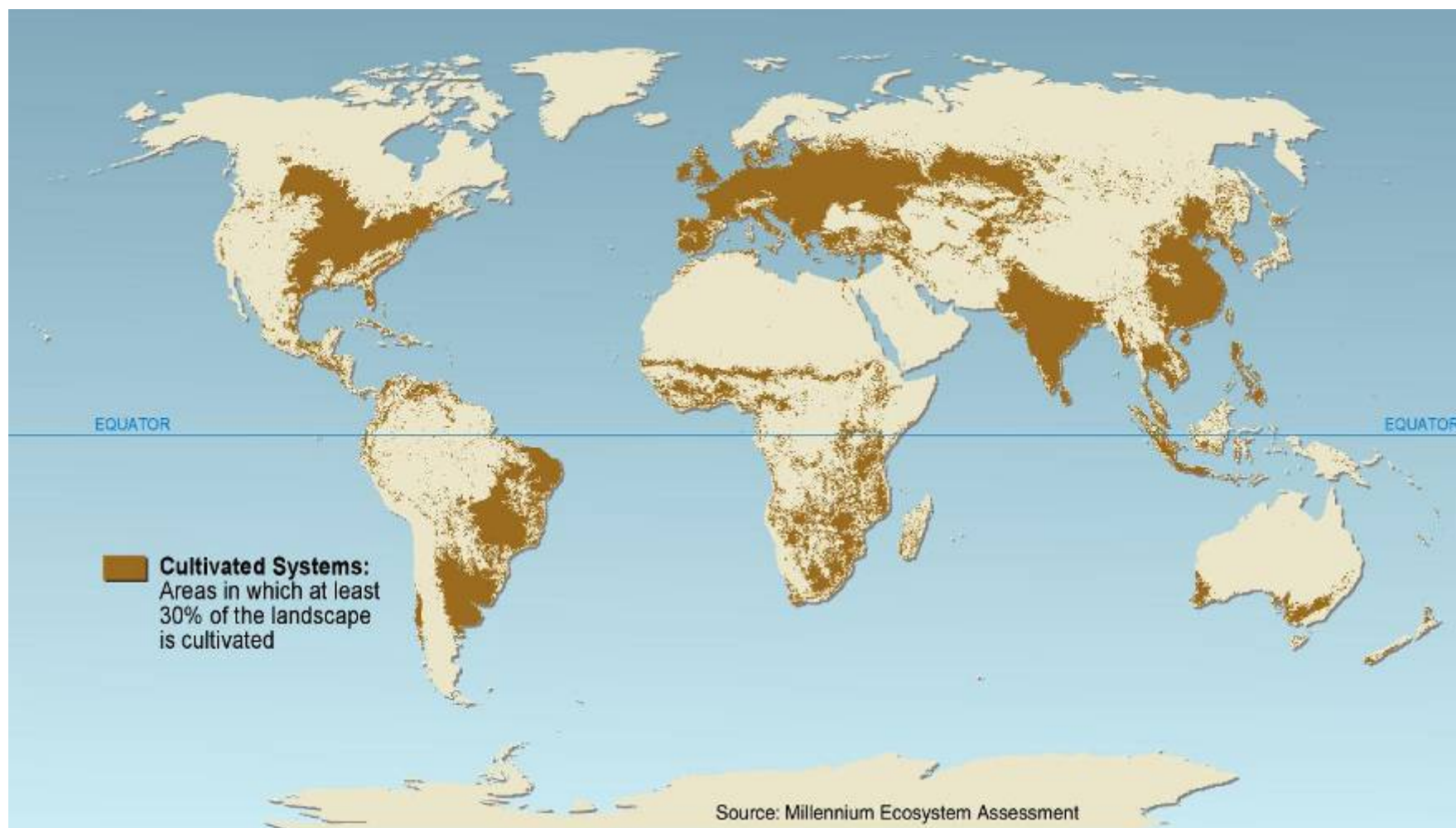
Figure 1 | Number of EID events per decade. EID events (defined as the temporal origin of an EID, represented by the original case or cluster of cases that represents a disease emerging in the human population—see Methods) are plotted with respect to **a**, pathogen type, **b**, transmission type, **c**, drug resistance and **d**, transmission mode (see keys for details).



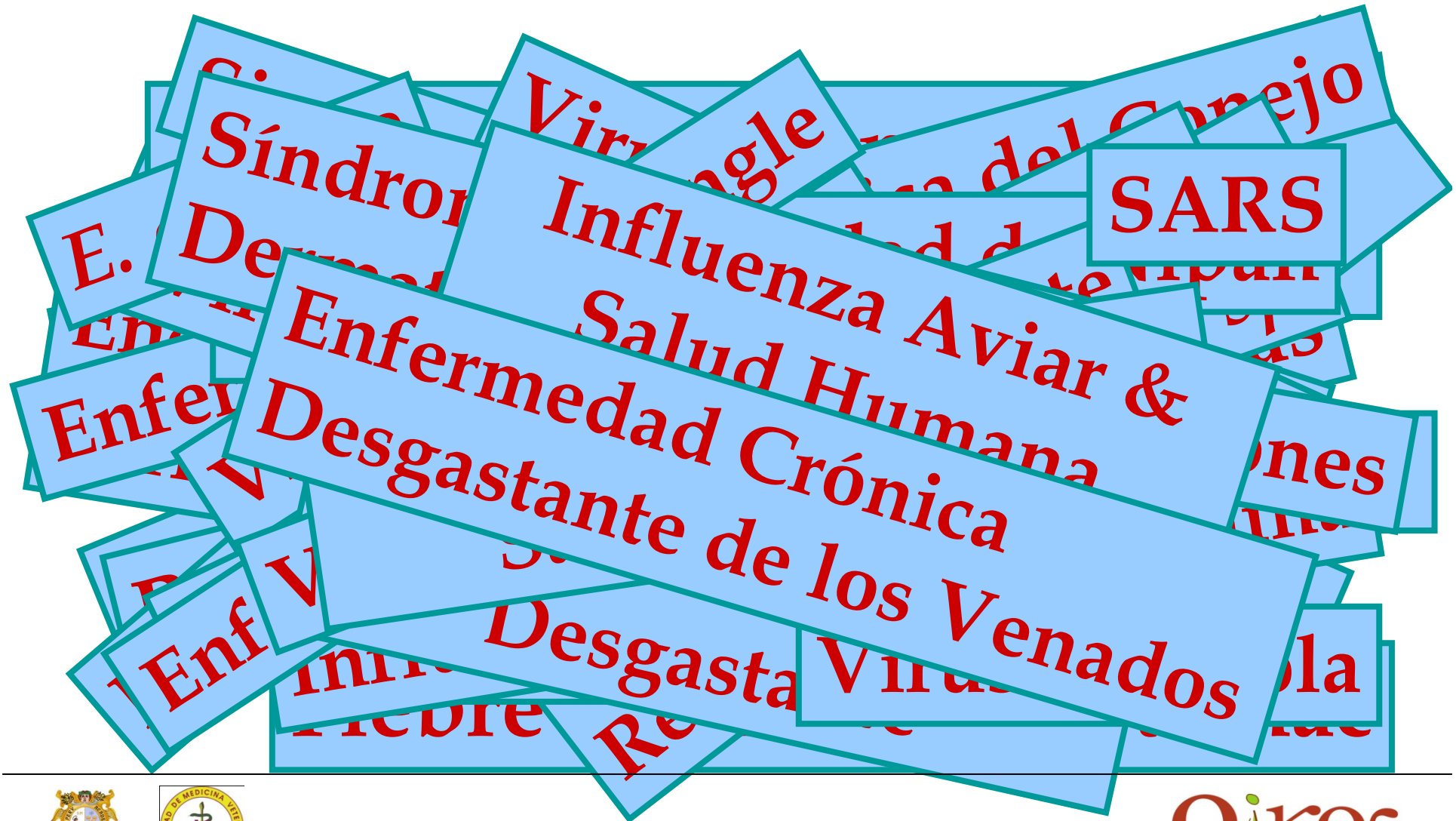


Source: NASA



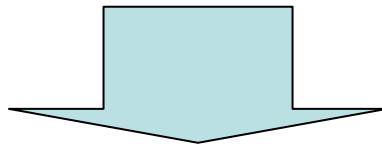


Nuevas Enfermedades



Factores que facilitan la dispersión de enfermedades emergentes

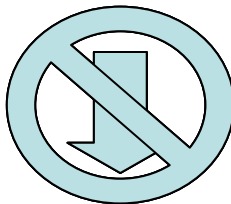
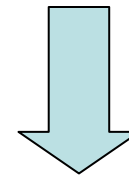
- Monoculturas y simplificación de habitat
- Destrucción de habitat
- Penetración humana en áreas “aisladas” (Invasión a nuevos nichos ecológicos)
- Decline de predadores
- Dominancia de generalistas sobre especialistas



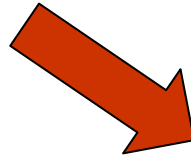
PÉRDIDA DE SERVICIOS ECOSISTÉMICOS



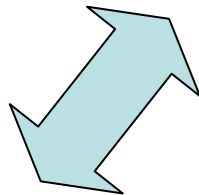
Hendra virus



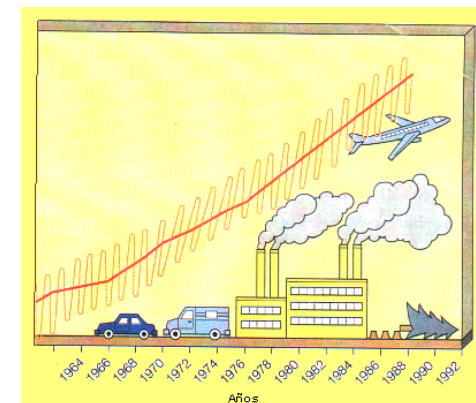
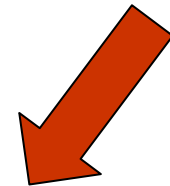
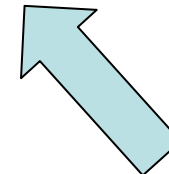
Relaciones
multifactoriales



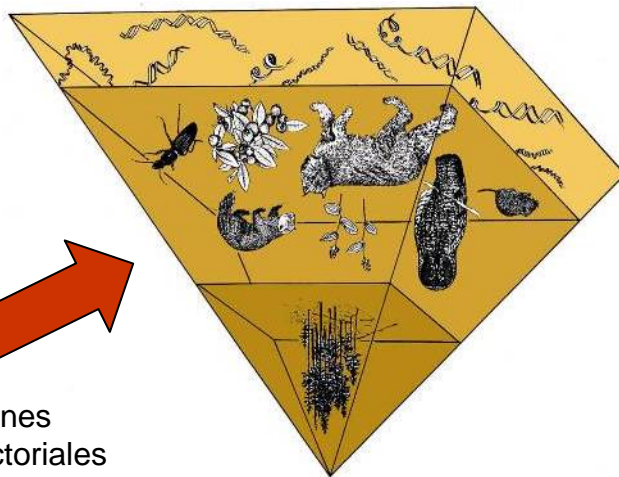
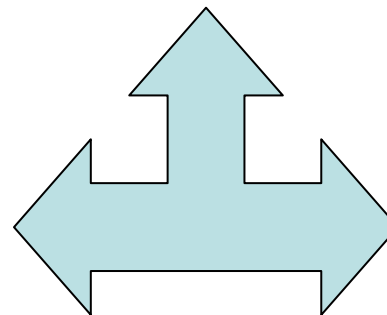
Enfermedades emergentes



Relaciones
multifactoriales

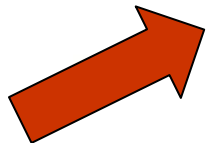


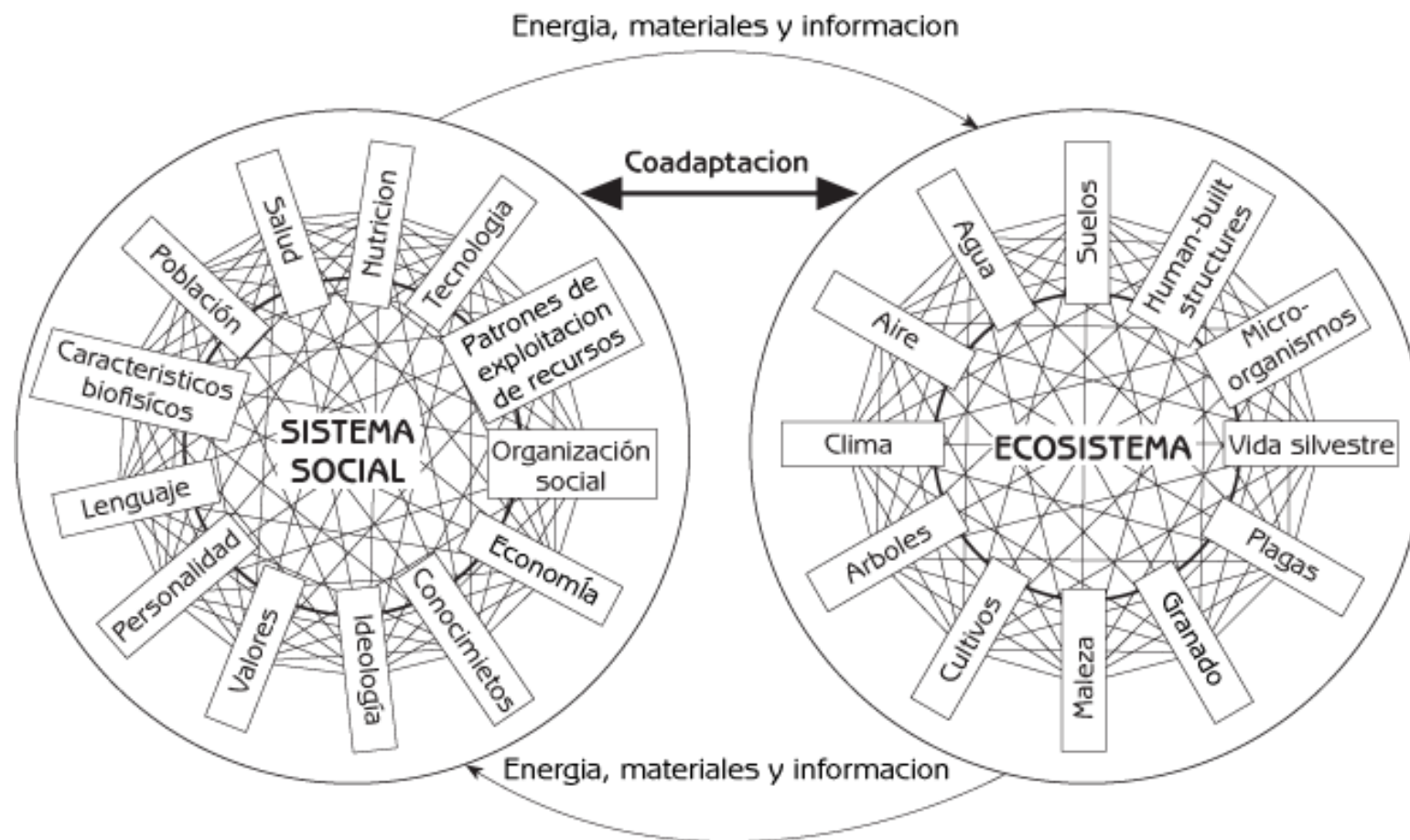
Cambio global



Pérdida de biodiversidad

Relaciones
multifactoriales

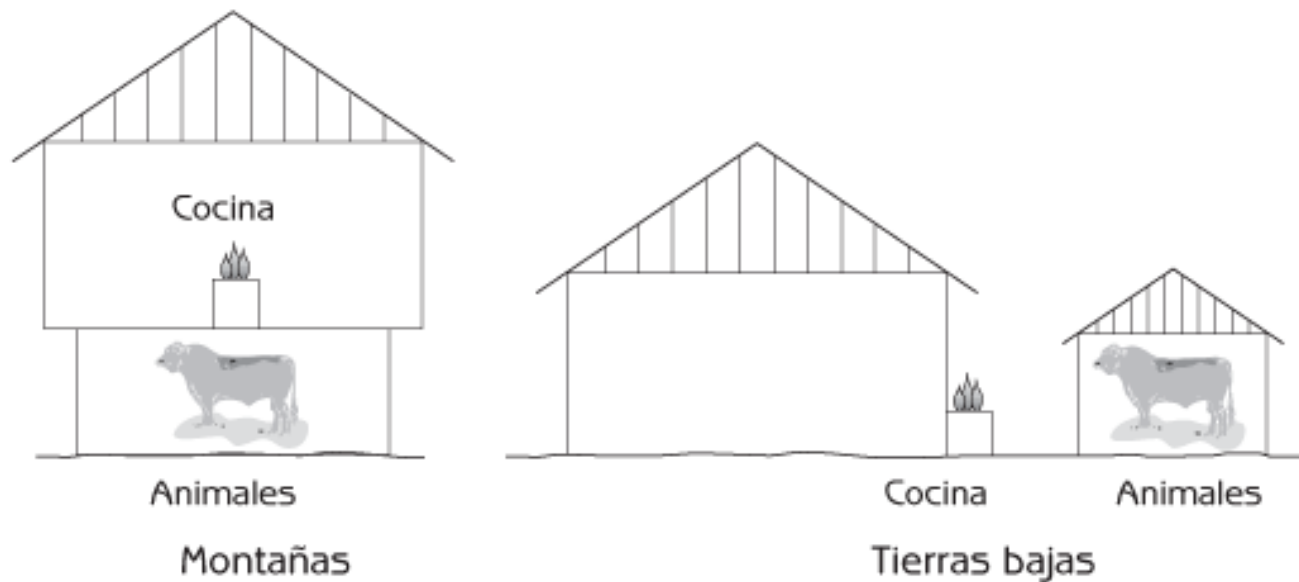




Interacción, coevolución y coadaptación del sistema social humano con el ecosistema. Fuente: Adaptado de Rambo, A. & Sjise, T. (1985) *An Introduction to Human Ecology Research on Agricultural Systems in Southeast Asia*, University of the Philippines, Los Banos, Philippines



Vivienda tradicional de las montañas y tierras bajas en Vietnam.



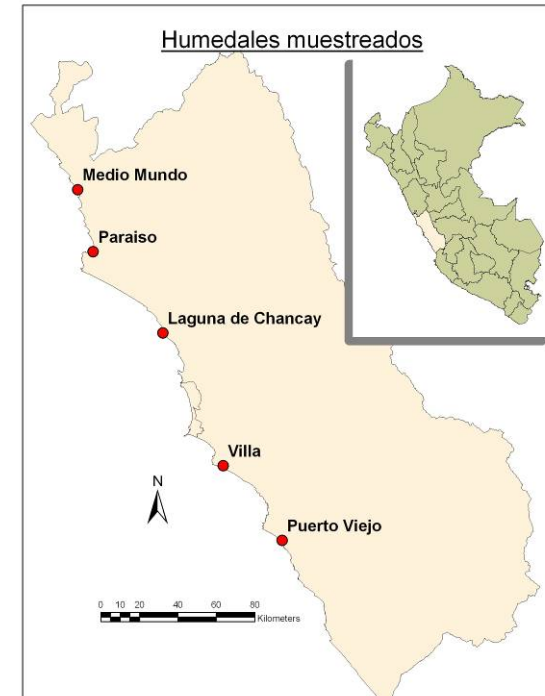
Las sociedades tradicionales típicamente están bien coadaptadas por tener lazos fuertes a su sistema de apoyo ambiental que les permiten conocer a fondo al mismo.





Avian influenza wild birds surveillance

- Determine the subtypes of AI circulating in wild birds in Peru
- Determine the bird species that act as carriers
- Five different wetlands in central Peru during 3 ½ years (2006-2009)
- 6887 environmental fecal samples collected from 34 avian species (13 families)
- 30 Avian Influenza isolates, 12 different subtypes.





AIV and NDV surveillance in Peru: An update from the 2006-2007-2008

Migratory season and H7N3 AIV strain isolation

Rosa I. Gonzalez¹, Eliana Icochea¹, Armando E. Gonzalez¹, Bruno M. Ghersi^{1,2}, David L. Blazes², Joel M. Montgomery²

¹ Universidad Nacional Mayor de San Marcos, Peru, ² US Naval Medical Research Center Detachment, Peru,



BACKGROUND

Migratory waterfowl are currently considered to be the primary reservoirs for avian influenza viruses. Although intense influenza virus surveillance in wild birds has been occurring in Europe, North America, Asia and Africa, few activities are present to date in South America. Environmental fecal samples were collected from four wetlands along the coast of central Peru from June 2008 to December 2008. Samples were processed for viral isolation in SPF embryonated chicken eggs. Allantoic fluids were evaluated for presence of hemmagglutinating agents (PMV and AI). A total of 4478 samples were collected during thirty months. Sixteen avian influenza isolates, representing eight different subtypes, and eight paramyxovirus isolates were obtained. Between the avian influenza isolates we report the presence of a low pathogenic H7N3 subtype from wild waterfowl (*Anas platyrhynchos*) in the following wetland.

METHODS

Study Sites

Four wetlands, along the central coast of Peru, were selected for sampling between June 2008 through December 2008 (Figure 1). These areas have the highest biodiversity of wild birds and are the largest wetlands in the department of Lima. All of them have contact with human habitation, poultry farms and/or pig farms.

Sampling locations



Figure 1.

Sampling Procedure

Bird colonies were identified and observed for 15 to 20 minutes by personnel experienced in bird identification. Only colonies conformed by one species were selected, or in case of more than one species been present colonies were identified as family. After the colony left the resting area, fresh fecal samples were immediately collected. Each fecal sample was collected with a sterile tipped applicator and placed in a cryovial containing viral transport media. Samples were then transported at 4°C to the laboratory to be processed within 24 hours of collection.

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1. Henny W. Avian Influenza. Field Manual of Wildlife Diseases. Madison, Wisconsin: US Geological Survey, Biological Resources Division, National Wildlife Health Center; 1996. p. 175-6.
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3. Chen RH, Shao HJ, Li XL, Zhou YJ, Hou YQ, Guo XJ, et al. Surveillance and identification of influenza A virus infection from migratory wild waterfowl in China (2004-2005). J Virol Methods [Int J Virol Methods]. 2008 Jan;143(1):65-70.
4. Hink A, Muller KJ, Wernke G, Glaser A, Kuehn E, Salazar A, et al. A serological survey in migrating waterfowl and other waterfowl in one of the most important resting sites of Germany. J Virol Methods [Int J Virol Methods]. 2008 Apr;143(1):101-10.
5. Parashar S, Smith N, Borth N, Brown V, Buck PA, Brunt A, et al. Wild bird influenza virus. 2005. Emerg Infect Dis. 2008 Jan;14(1):146-7.
6. Rappelli B, Hink A. 2. Birds and influenza H5N1 virus movement in and within North America. Emerg Infect Dis. 2008 Oct;14(10):1499-92.

Positive Samples

Date	Place	Common Name	Virus	HN
16-Aug-06	Puerto Viejo	Ruddy Turnstone	PMV	
16-Aug-06	Puerto Viejo	Cormorant	PMV	
16-Aug-06	Puerto Viejo	White-checked pintail + Cinnamon Teal	PMV	
17-Oct-06	Puerto Viejo	Ruddy Turnstone	AI	H10N9
24-Oct-06	Puerto Viejo	Ruddy Turnstone	AI	H10N9
6-Nov-06	Puerto Viejo	White-checked pintail + Cinnamon Teal	AI	H1N8
7-Nov-06	Puerto Viejo	American Oystercatcher	AI	H10N9
8-Feb-07	Medio Mundo	White-checked pintail + Cinnamon Teal	AI	H4N3
13-Feb-07	Medio Mundo	White-checked pintail + Cinnamon Teal	AI	H4N3
13-Feb-07	Medio Mundo	Parusian Pelican	AI	H4N3
6-Jun-07	Paraiso	Duck	PMV	
6-Jun-07	Paraiso	Gull	PMV	
11-Jun-07	Paraiso	Egret	PMV	
30-Oct-07	Villa	Duck	PMV	
30-Oct-07	Villa	Egret	PMV	
20-Nov-07	Paraiso	Willet	AI	H1N2
20-Nov-07	Paraiso	Dominican Gull	AI	H1N2
22-Apr-08	Puerto Viejo	Ruddy Turnstone	AI	H1N2
22-Apr-08	Puerto Viejo	Ruddy Turnstone	AI	H1N2
22-Apr-08	Puerto Viejo	Common Goldeneye	AI	H1N2
17-Jul-08	Villa	White-checked pintail + Cinnamon Teal	AI	H7N3
22-Jul-08	Villa	White-checked pintail + Cinnamon Teal	AI	H2N9
29-Oct-08	Medio Mundo	Ruddy Turnstone	AI	H10N7
5-Nov-08	Medio Mundo	American Oystercatcher	AI	H10N7

H7 Cleavage site evaluation

Avian Paramyxovirus (H7N3)	E I P K G R - - - G L F	LP
Avian Paramyxovirus (H2N9)	E I P K G R - - - G L F	LP
Avian Paramyxovirus (H5N1)	E N P K R T - - - G L F	LP
Human Paramyxovirus (H5N1)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N1)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N2)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N3)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N4)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N5)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N6)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N7)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N8)	E I P K G R - - - G L F	LP
Human Paramyxovirus (H2N9)	E I P K G R - - - G L F	LP



CONCLUSIONS & DISCUSSION

Our findings provide additional evidence that AI viruses are present and circulate in wild birds in South America, and thus can be potential sources of influenza viruses perhaps spreading infection among a variety of avian species, including poultry, as well as swine and humans, along their flyways between North and South America.

Even though all isolates are from LP strains, the presence of an H7N3 represents a threat.

1. **DISCLAIMER:** The views and opinions of the authors are those of the authors and do not necessarily reflect the official policy or position of the Department of the Navy, Department of Defense, nor the U.S. Government.
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EVALUACIÓN SANITARIA DE ANIMALES DOMÉSTICOS, SILVESTRES Y HUMANOS EN LAS COMUNIDADES DE NUEVA ESPERANZA (RÍO YAVARÍ-MIRIM) Y SOL NACIENTE (RIO AMAZONAS)

OBJETIVOS

- Estudio de la prevalencia de enfermedades de mamíferos domésticos y mamíferos silvestres cazados.
- Evaluación sanitaria de los animales domésticos de abasto en las comunidades.
- Evaluación de la prevalencia de la malaria en las poblaciones humanas.
- Desarrollo de un programa de educación para las comunidades locales enfocado en la conservación, cría de animales domésticos y aspectos ecológicos sanitarios de los animales y de las poblaciones humanas.



INSTITUCIONES

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EVALUACIÓN SANITARIA DE ANIMALES DOMÉSTICOS, SILVESTRES Y HUMANOS EN LAS COMUNIDADES DE NUEVA ESPERANZA (RÍO YAVARÍ-MIRIM) Y SOL NACIENTE (RIO AMAZONAS)

Análisis de Malaria

Nueva Esperanza

Plasmodium Falciparum	6
Plasmodium Vivax	15
TOTAL INFECTADOS	21
TOTAL PERSONAS	78
PREVALENCIA	26,9

Prevalencia de Malaria en función de la presencia de sintomatología clínica (n=78)

SX	TOTAL	NEGATIVOS	POSITIVOS	FALCIPARUM	VIVAX	% POSITIVOS	% FALCIPARUM	% VIVAX
CON SX	12	8	4	1	3	33,3	8,3	25,0
SIN SX	65	48	17	5	12	26,2	7,7	18,5

Prevalencia de Malaria en función del trabajo de madera (n=78)

MADERA	TOTAL	NEGATIVOS	POSITIVOS	FALCIPARUM	VIVAX	% POSITIVOS	% FALCIPARUM	% VIVAX
SI MADERA	32	22	10	3	7	31,3	9,4	21,9
NO MADERA	45	34	11	3	8	24,4	6,7	17,8
TOTAL	77	56	21	6	15	27,3	7,8	19,5

Necesidades de investigación

- Integración entre la salud humana, animal y ecosistémica.
- Interacción entre los procesos ecológicos y evolutivos para determinar las dinámicas de las enfermedades.
- Rol de la ecología entre hospederos, patógenos y las interacciones entre múltiples patógenos.
- Medición del riesgo y vigilancia de la emergencia de enfermedades a través de modelos predictivos teniendo en cuenta factores medioambientales.
- Monitorear la salud de los animales domésticos y silvestres



Acciones necesarias

- Identificación de 'hotspots' de potenciales enfermedades emergentes.
- Vigilancia de la vida silvestre.
- Desarrollo de estrategias de mitigación de enfermedades emergentes.
- Mantener la biodiversidad y los servicios que nos ofrece.



Acciones necesarias

- Creación de redes de trabajo interdisciplinarias e interinstitucionales.
- Mayor intercambio de información.
- Utilización de los principios de la ecología para el control de las enfermedades de hospederos múltiples, focalizándose en el descubrimiento y caracterización de patógenos.
- Desarrollo de vacunas.





La salud de todas las especies está conectada a través de las realidades ecológicas que gobiernan la vida en el planeta.



MEDICINA DE LA CONSERVACIÓN

Aproximación transversal al término de SALUD

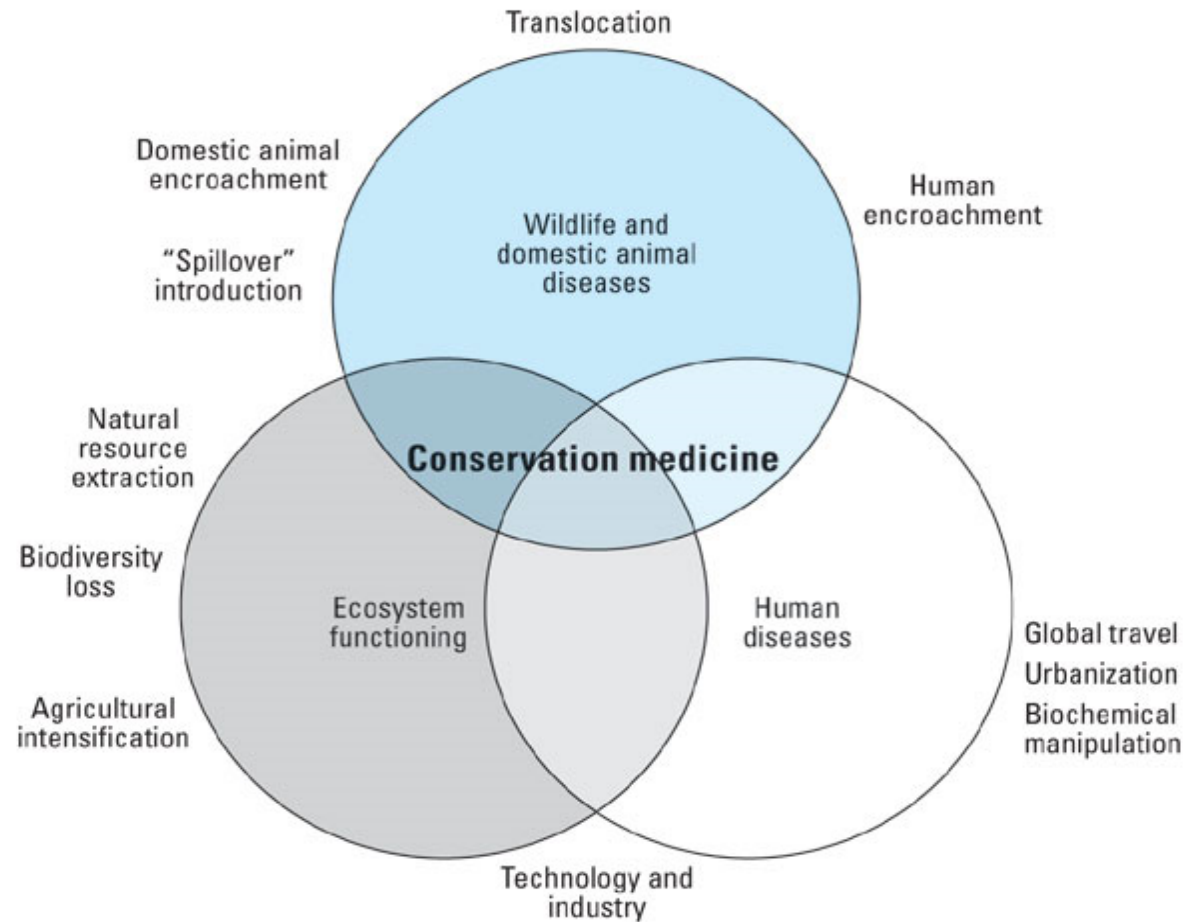


Figure 2. The main elements converging under the Consortium for Conservation Medicine. Conservation medicine combines conservation biology, wildlife veterinary medicine, and public health. Adapted from Tabor (2002).



**MUCHAS GRACIAS POR
SU ATENCIÓN**



“Servicios ecosistémicos, ecología de las enfermedades y bienestar humano”

Eric Chávez Betancourt

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